Quick Reference Guide - Using WebBLAST

# To Use WebBLAST (see detailed instructions below):

Before you use WebBLAST for the first time, the Site Administrator will need to provide you with a User Name and Password, and assign you to a Project and Group. You must install the Sequence Uploader on your computer (for installation instructions, see <a href="http://research.nhgri.nih.gov/webblast/client/">http://research.nhgri.nih.gov/webblast/client/</a>). For additional assistance, NHGRI users can contact Lan Support.

- 1. Login.
- **2.** Change Password (if this is the first login).
- 3. Add a subproject.
- 4. Associate group with subproject.
- 5. Setup a subproject.
- **6.** Upload sequence.
- 7. Select a project and subproject.
- 8. Check sequence and BLAST status.
- **9.** Retrieve data.
- 10. ReBLAST as needed.

\*Note: The Admin link and associated functions are available only to users who have the Administrator or Site-Admin privilege.

#### Login:

- Enter the URL for the WebBLAST CGI in your web browser (for NHGRI users, enter: http://hercules2.nhgri.nih.gov/webblast).
- 2. Enter your User Name and Password.
- 3. Select Login.

# To Add a Subproject:

- 1. Login to WebBLAST.
- 2. Select the Admin link.
- 3. Select Add Subproject from the Administration Menu.
- 4. Enter the Subproject Name.
- Select a Project to associate the subproject to from the drop-down list.
- 6. Select Add Subproject.

### To Setup a Subproject:

- 1. Login to WebBLAST.
- 2. Select the Admin link.
- 3. Select Setup Subproject from the Administration Menu.
- 4. Select the Project :: Subproject from the drop-down list.
- 5. To setup BLAST and filters, select BLAST & Filters from the Action drop-down list. Note: WebBLAST requires a BLAST configuration in order to run. At minimum, you must select a BLAST program and database. If the "month" database is selected, all sequences will be requeued for BLASTing every 30 days.
- 6. To setup masks, select Masks from the Action drop-down list.
- 7. Select Setup Subproject.
- **8.** Enter the data in the fields as necessary.
- 9. For BLAST and Filters, select Save Configuration.
- 10. For masks, select Add to Configuration.
- 11. To remove a configuration, place a check in the check box to the left of the configuration and select Remove Selected Configuration.

\*Note: You can define as many BLASTs & Filters and Masks as you desire, as long as you use different combinations of BLAST programs and databases.

# Change Password:

- 1. Login to WebBLAST.
- 2. Select the Change Password link.
- **3.** Enter the User Name, Current Password, New Password, and Retype New Password.
- 4. Select Change Password.

#### To Associate Groups with Subprojects:

- 1. Login to WebBLAST.
- 2. Select the Admin link.
- 3. Select Manage Groups from the Administration Menu.
- 4. Select the appropriate Project :: Group from the drop-down list.
- **5.** Select Associate Group with Subprojects from the Action drop-down list.
- 6. Select Manage Group.
- 7. Highlight the appropriate subprojects that you wish to associate with the group from the Subprojects list and move them to the "Group Name" Group list by using the right arrow.

# To Upload Sequences (using the Sequence Uploader):

- 1. Open the Sequence Uploader program.
- 2. Select Add File.
- 3. Select the sequence or directory and select Open.
- 4. Select Submit.
- 5. Enter your WebBLAST User Name and Password on the Login screen.
- 6. Select OK.
- 7. Select the appropriate project from the drop-down list and select
- 8. Select the appropriate subproject from the drop-down list and select OK.
- 9. Select Quit to end the Sequence Uploader session.

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### Select a Project and Subproject:

- 1. Login to WebBLAST.
- 2. Select a Project from the drop-down list.
- 3. Select Enter Project.
- 4. Select a Subproject from the drop-down list, and/or
- 5. Enter a Sample Name (entering both the Subproject and Sample Name filters the search for data).

#### Retrieving Data:

- 1. Login to WebBLAST.
- 2. Select a Project.
- 3. Select a Subproject and/or enter a Sample Name.
- 4. Select Retrieve Data.
- 5. To review the results for a particular sample, select the "sample name" link.

#### To ReBLAST all Sequences within a Subproject:

\*Note: ReBLASTing all sequences is computationally intensive. Ensure that you want to reBLAST all sequences before performing this step.

- 1. Login to WebBLAST.
- 2. Select the Admin link.
- 3. Select Setup ReBLASTs from the Administration Menu.
- **4.** Select a Project :: Subproject to reBLAST from the drop-down menu.
- Deselect the Select Individual Sequences check box, if already checked.
- 6. Select the reBLAST button.
- \*Note: If the "month" database is selected, all sequences will be requeued for BLASTing every 30 days.

# To Review Retired BLASTs and Sequences:

- 1. Login to WebBLAST.
- 2. Select a Project and a Subproject and/or enter a Sample Name.
- 3. Select Retrieve Data.
- 4. Select the "sample name" link for the results you wish to view.
- 5. Select the View Retired BLAST link or the View Retired Sequence link (the links are available only if a retired BLAST or retired sequence exists).
- Select the link of the retired BLAST or sequence you wish to view.

#### To Save the SCF or ABI Trace:

- 1. Login to WebBLAST.
- 2. Select a Project.
- 3. Select a Subproject and/or enter a Sample Name.
- 4. Select Retrieve Data.
- 5. Select the "sample name" link for the results you wish to view.
- **6.** Select the Save SCF Trace button or the Save ABI Trace button.

#### Windows:

- 7. Select the Save this File to Disk button. Select OK.
- 8. Select the file directory in which you wish to save the trace by using the Save in: drop-down list.
- **9.** Enter the name of the trace file and select Save.

#### Mac:

- 11. Select Save File.
- 12. Select the file directory where you wish to save the trace file.
- 13. Enter the name of the file in the Save this document as: text box and select Save.

# Checking Sequence and BLAST Status:

- 1. Login to WebBLAST.
- 2. Select a Project.
- 3. Select a Subproject and/or enter a Sample Name.
- 4. Select Check Status.
- **5.** To display the results for a particular sample, select the associated "sample name" link.

*Note:* For the "sample name" link to be active, the BLAST status must be "finished".

# To ReBLAST Individual Sequences Within a Subproject:

- 1. Login to WebBLAST.
- 2. Select the Admin link.
- 3. Select Setup ReBLASTs from the Administration Menu.
- **4.** Select a Project :: Subproject to reBLAST from the drop-down menu.
- 5. Check the Select Individual Sequences check box, if not checked already.
- 6. Select the reBLAST button.
- 7. Highlight the sequences you wish to reBLAST.
- 8. Select ReBLAST.
- \*Note: If the "month" database is selected, all sequences will be requeued for BLASTing every 30 days.

# To View BLAST Reports:

- 1. Login to WebBLAST.
- 2. Select a Project and a Subproject and/or enter a Sample Name.
- 3. Select Retrieve Data.
- 4. Select the "sample name" link for the results you wish to view.
- **5.** Select the sample\_name.database.BLAST\_program link to see the entire BLAST report.
- 6. To access the BLAST alignment for a specific sequence, select the "sequence name" link on the Results for 'Sample Name' screen.

# Launching the Trace Viewer:

*Note:* The Launch Trace Viewer button is only displayed if a trace was uploaded with the sequence data.

- 1. Login to WebBLAST.
- 2. Select a Project.
- 3. Select a Subproject and/or enter a Sample Name.
- 4. Select Retrieve Data.
- 5. Select the "sample name" link for the results you wish to view.
- 6. Select Launch Trace Viewer.
- 7. To view the number of data points, select the Data button.
- **8.** To close the Trace Viewer, select the Hide or Close button.
- **9.** If you adjusted the view on the Trace Viewer, select the Reset button to return to the default display.

# To Save the Sequence Data Locally:

- 1. Login to WebBLAST.
- 2. Select a Project.
- 3. Select a Subproject and/or enter a Sample Name.
- 4. Select Retrieve Data.
- 5. Select the "sample name" link for the results you wish to view.
- 6. Select the Save Sequence or Save Masked Sequence button.
- 7. Follow the dialogs for Windows and Mac to save the sequence to a file directory.